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Fig. 1

EGP-2 promoter sequences

-3967. ^{BglII}AGATCTAGAA TAGAGAGGGA TTGCTGTCAT AGTGGTTAAG GACTTTTACT CTTCAATCTA TATAAAGGAC TTTTGTTCCTTC
 -3987. TACTCATCTA TTACTTATGG GATAACAAA ATTTTATAGAA CTGGTAGTCT AATTTATAT ATATATATAT ATATATATAT
 -3987. ATATATATAT ATATATATAT ATATATTTT TTTTATTTT TTTTAGACAG AGTTTGTCTC TTGTTGCCCA GGCTGGAGTG
 -3727. CAATGGCATG ATCTTCGCTC ACCACAACCT CCGCTCCTCG GGTTCAAAGTG ATTCTCCTGC CTCAGCCTCC CAAGTATCTG
 -3647. GAATTCACAG CATGTGCCAC CATGCCACG TAATTTTAT ATTTTATAGTA GAGACAGGTT TTCACACAGGT TGCCACAGGCT
 -3567. GCTCTCAAC TCCTGACCTC AAGTGATCCA CCGCTTTGG AATATGGCAC TAGAACTACA AAAGATTCAC. AATTAACACA TAAACCGAGT
 -3487. CTAGCCCTGAA AATATTAATA AATGTGCTTA AATATGGCAC TAGAACTACA AAAGATTCAC. AATTAACACA TAAACCGAGT
 -3407. AATTTTGAGC AAAGATGAC AAATTGAGAA GGTTTAAATG AGTAACTAAA ATAAACAATA CCGGCGG (GTGCAGTGGCTCA p39^B
 -3327. TGCCTGTAAT CCCAGCACTT TGGGAAGCTG ^{SpI}AGGCGGGTGG ATCACCCTGAG GTCAGGAGTT CAAGACCAGC CTGGCCCAACG
 -3247. TAGTGAACCC CGGTCTCTAC TAAAAATACA AAAATTAGCC GGCGGAGTG GCAGGCGCT. GTAATCACAG CTAATCGGGA
 -3167. GGCTGAGACA GGAGATTTGC TTGAACCCAG GAGGTGAGG TTGCAGTGAG CTGAGAACAC GCCATTGTAC TCCAGCCTGG
 -3087. GTAAACAGAT TGAACACTTA TCTTAAAAA AAAAAAAGG CGGACACGGT GGCTTGACCC TGTATCCCA GCACCTTGGG
 -3007. AGGCCGAGGC AAGAGGATCA CAAAGTCAGG AGATCAGAC CATCTGGCC AACATGTGTA AACTCTGTCT CAACTGNAAA
 -2927. TACAAAATTT AGCCGGTGT GTTGGTGGG GCCTGTATC CCAGCTATTC AGGAGGCTGA GGCAGGAGAA TTGCTTGAAC
 -2847. CCAAGAGGTG GAGGTTGCAG TCCGCCAAGA TCATGCCACT GCACCTGACG TTGGTGACCA GAGCAAGACC CCATCTCAAA
 -2767. AAAAAAATAA AAGAAAAAAT ACCCTGGATC AGCCGGGTGT GTTGGCTCAA GCCTGTATC CCIAGCACTTTGGAGGCTGA p39^{B4-7}
 -2687. GGTGGGCAGA TCACCTGAGG TCAGGAGTTC AAGACCAGCC TGACCAACAT GGAGAAACCC CATCTCTACT AAAAATACAA
 -2607. AAAATTAGCC GGACGTGGTG GCACATGCTT GTAATCCAG CTACTCAGGA GGCTGAGGCA GGAGATTTGC CTGAATCCGG
 -2547. GAGGCGGAGG TTGTGGTGAG GTGAGATGAT GCCATTGCAC TCCAGCCTGG GCAACACAG CAAAACTCTG CCTCAAAAAA
 -2447. AGAAAGAAA AAAAAAAGA AAAAAAGAA AAAATACCT GGATGTATAC TCAGATACAA ^{AP-1}TGAGTCAGAG ATTAGTCTGG
 -2367. TATTTTGTCA TTTATTTAAT AATTATGCTT ACTCAATCA CTT (TATTGTAAATTAACAATA AATAGCTGC CAGTTATAAG p39^{B17-1}
 -2287. AAGATGAAGT TCTCCGATTT AGTAAACAG AATTAGACCT CAGAAATGGAA CATTTTGCCA ATAAAGCCAC AATAACCACT
 -2207. TAGTTTATTC TTGGGAAAG TATATGTAAT TTGGAGAAAG GCAAACTTCC ^{Sp-1}GAAACATC CAAATTCAG CAGACAACAA
 -2127. AAATCTGGTT AACTTCTTCC ^{Sp-1}GATTGTGA GTACTATTC (TTTTTTTTTTT TTTGTTTGT TTTTTTTTTT GAGACGGAGT p39^{B15-1}
 -2047. TTCGCTCTTG TTGCCCAGGC TGGAGTGCAA TGGCGAATG TTGTTCACT GCAACCTCTG CCTCCAGGT TCAAGTGAAT
 -1967. CTCCTGCCCTC AGTCTCCTGA GTAGCTGGGA TTACAGGCC ^{Sp-1}CCGACCCAC GCCTGGCTAA CTTCTGTAT TTTTAGTAGA
 -1887. GACGGGGTTT CACCATGTTG GCCAGGCTGG TCTCGAATC CTGACCTTAG GTGATCCCGC ^{Sp-1}CGCTCGGCC TCCCAAGTG
 -1807. CTGAGATTAC AGGCATGAGC CACCGTACCT GGCCTAATA CCTATTTC TATACCACGT GAAATTTAAA TTATACAAAA

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Fig. 1, contd.

-1727. CAATATATAG AGGTACTTAG AACAGCATGA CTAATTACAT TAATCAACTT GCCGGCACTT CAACAGAATA CAACATAGAA
-1647. ATGATTGTTT TAATATAAC ATAAGCTTTG AATTGACATA TACTTGAGAA AATTAATCAA ACTTAGCTGA ATCTTAAAT
-1567. TGCCTTTTTC CTTTCTCTCT TTTTCTTTTA TTTTCTTATT TTTTGAGANG GAGTCTTGCT CTGTTGCCAG ACTGGAGTGC
-1487. AGCGGTTTGG TCTCGGCTCA CCGCAACCTC CGACTCTCTG GTTCAACGGA TTCTCCTGCC TCAGCCTCCT GAGTAGCTGG
-1407. GATTACAGGT GCCTGCCACC ACACCTGGCT ACCTTTTGTG TTTTCTAGTTG AGATGGGTTT CACCATGTTG GCCAGGATGG
-1327. TCTCGAATC CTGACCTCGG ATCTGCCAC ATCTGCCAC CTGTGCCCC AGCAAGTGC TGGATTACA AGCATGAGCC ACCGTGCCCA
-1247. GCCTCCTTTC CTCTTTTAA CTCTTACTT TATGATTTCT TTAGTGGATA AAAAGCTTTT AAAAATAGG TTACATGAT
-1167. ATTACAGCTA ACAAAAATA ACATTAAAA AACTAAATA GTATATATAT GAATATTTA TAATTATTTT AATATTGTAA
-1087. TAATATAGT TGTGTGAT TGAATTCATC TGCACGGAA TCGATTACTG TCCTTTCTTT CTATTTCCCT ATATTTTCTT p39¹⁷⁻²
-1017. TCCGAGGCT CATCAACAT TTGGTTCTTT AATAGTAACC AAAACCGAA ATCATCTCG TTCTCAGTAT TTGGCTCTAT
-937. GCGAAGCTT TTTCTTTTCT CTCTTTTCTT TTTTCTTGA GACGGAGTCT TGCTCCTGTC GCCCAGGCTG GAGTGTATG
-857. GCACGATCTC TGCTCAGTC AACCTCAGCC TCCCCAGTAG CTGGGATTAC AGCATGCGC CACCACGCCG GGCTAATTTT
-777. GTATCTTTTA GTAGAGCGG CTTCTCTCA TGTGTGTCAG GCTGTCTCG AACTTCAAC CTCAGGTGAT CCGCGGCT
-697. CGGCTCCC [AAGTGCTAG ATTACAGCG TGAGCCACCG CGCTACCTT GGAACACCT TTTCTTACAT CTTCAGTGC p39²⁴⁻¹
-617. TAGAATGCT TAGAATAACG AAAAAGAT TATTAAGAT AATTATAAG AACACTCAT TTTCTTCCCA AGAGGCCAA
-537. GATTCTTCT TTTCTTTTCT TTTCTTTCTA ATTTCAAGG AGTATAATTA AATTGCCAGG TAAAGCTGA
-457. AGCTTTTCTT TTATAGTGT CTGGAAGTT CTCTGCTGT GTTGTATTT CTTTACCTT CCACCTTCTT CTATCCAGTT
-377. CCGCACCTT TCCCCCAGG CCCCATTTCT CAAGCTTTCAGAGCAGGCT CTTCCGTTA [AAGGAGTCT TCAGCACAGA p39⁵¹¹⁻¹
-297. ATCTTCAAC CTCTCTGGAG GCCACCAAG ATCCCTAAC CCGCCTAGA GACGAAGCAC CTGAGGCGG GCGGAGCGGG
-217. GCGGCGGGC CCACACCTCT GAGAGGGCC GCGCCCCAAC TGCAGCGCGG GGGTGGGG AGGGAGGCT ACTCACTCCC
-137. CCAACTCCG GCGGAGTCT CATTACAGAG CACACAGCGG CAGAGGTGAG CAGTCCCGG AAGGGGCGA GAGGCGGFC
-57. [CGCAGGTCCGAGGTGT CGCTCCGCTT CCGGCGCGC ACAGAGGCT AGTCTTCTG CGAGCAGACCTTCGAGCC p39^{512-2,3}
+23. GGTCCGGGA CCCCCTGTC GCTGTCTCC CGAGCGGAC CCGCTGCCC CAGGCTCTCTT CCGCGCGCG GCAGCATGCG
+103. TGTCCACTC CCGCGCAG CCGTCCCGG AGTCCCGG CCGTCCCGG CCGCTCTCTT CCGCGCGCG GCAGCATGCG
+183. GCGGCGGAG GTCTGCGCTT TGGGCTTCT GTTTCGCGG GCGACCGCA CTTTTCGCG AGCTTCTGAA GGTGAGGCGC
+263. GGAATGAGC AGATTGTCT AGCTGGGCTG GCGTGGGG CA

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Fig. 2

